

Comparison with single nuclei RNA-seq (sNuc-Seq) data

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An abbreviated version of this protocol was published in eLIFE in Oct 2018

Species and cell-type properties of classically defined human and rodent neurons and glia

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Detailed protocol

All code and outputs for the comparative analysis can be found at: https://xu-xiao.github.io/non_transgenic_cell_type_profiling/html/figure_3_s3.nb.html

How to cite: (Readers should cite both the Bio-protocol preprint and the original research article where this protocol was used)

1. Xu, X. (2021). Comparison with single nuclei RNA-seq (sNuc-Seq) data. Bio-protocol Preprint. bio-protocol.org/prep1286.
2. Xu, X., Stoyanova, E. I., Lemiesz, A. E., Xing, J., Mash, D. C. and Heintz, N. (2018). Species and cell-type properties of classically defined human and rodent neurons and glia. eLIFE. DOI: [10.7554/eLife.37551](https://doi.org/10.7554/eLife.37551)

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